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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/780,438C

DATE: 05/08/2003

TIME: 14:42:57

Input Set : A:\EP.txt

Output Set: N:\CRF4\05082003\I780438C.raw

3 <110> APPLICANT: Qi, Xiaoyang  
 5 <120> TITLE OF INVENTION: Fusogenic Properties of Saposin C and Related Proteins and Polypeptides  
 6 for Application to Transmembrane Drug Delivery Systems  
 8 <130> FILE REFERENCE: 10872/0474352  
 10 <140> CURRENT APPLICATION NUMBER: US 09/780,438C  
 11 <141> CURRENT FILING DATE: 2000-02-11  
 13 <150> PRIOR APPLICATION NUMBER: US 60/181,754  
 14 <151> PRIOR FILING DATE: 2000-02-11  
 16 <160> NUMBER OF SEQ ID NOS: 6  
 18 <170> SOFTWARE: PatentIn version 3.1  
 20 <210> SEQ ID NO: 1  
 21 <211> LENGTH: 40  
 22 <212> TYPE: PRT  
 23 <213> ORGANISM: Homo sapiens  
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 28 1 5 10 15  
 31 Lys Leu Ile Asp Asn Asn Lys Thr Glu Lys Glu Ile Leu Asp Ala Phe  
 32 20 25 30  
 35 Asp Lys Met Cys Ser Lys Leu Pro  
 36 35 40  
 39 <210> SEQ ID NO: 2  
 40 <211> LENGTH: 38  
 41 <212> TYPE: PRT  
 42 <213> ORGANISM: Homo sapiens  
 44 <400> SEQUENCE: 2  
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 50 Ile Asp Asn Asn Lys Thr Glu Lys Glu Ile Leu Asp Ala Phe Asp Lys  
 51 20 25 30  
 54 Met Cys Ser Lys Leu Pro  
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 58 <210> SEQ ID NO: 3  
 59 <211> LENGTH: 38  
 60 <212> TYPE: PRT  
 61 <213> ORGANISM: Homo sapiens  
 63 <220> FEATURE:  
 64 <221> NAME/KEY: MISC\_FEATURE  
 65 <222> LOCATION: (1)..(1)  
 66 <223> OTHER INFORMATION: Where the amino acid located at 1 is a hydrophobic amino acids, i  
 67 ncluding Val, Leu, Ile, Met, Pro, Phe, and Ala  
 70 <220> FEATURE:

71 <221> NAME/KEY: MISC\_FEATURE ,

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72 <222> LOCATION: (2)..(2)
73 <223> OTHER INFORMATION: Where the amino acid located at 2 is an uncharged polar amino
aci
74      d, including Thr, Ser, Tyr, Gly, Gln, and Asn
77 <220> FEATURE:
78 <221> NAME/KEY: MISC_FEATURE
79 <222> LOCATION: (5)..(5)
80 <223> OTHER INFORMATION: Where the amino acid located at 5 is a hydrophobic amino
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81      cluding Val, Leu, Ile, Met, Pro, Phe, and Ala
84 <220> FEATURE:
85 <221> NAME/KEY: MISC_FEATURE
86 <222> LOCATION: (8)..(10)
87 <223> OTHER INFORMATION: Where the amino acids located at 8-10 are hydrophobic amino
acids
88      , including Val, Leu, Ile, Met, Pro, Phe, and Ala
91 <220> FEATURE:
92 <221> NAME/KEY: MISC_FEATURE
93 <222> LOCATION: (13)..(13)
94 <223> OTHER INFORMATION: Where the amino acid located at 13 is a hydrophobic amino
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95      ncluding Val, Leu, Ile, Met, Pro, Phe, and Ala
98 <220> FEATURE:
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100 <222> LOCATION: (14)..(14)
101 <223> OTHER INFORMATION: Where the amino acid located at 14 is an uncharged polar
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102      id, including Thr, Ser, Tyr, Gly, Gln, and Asn
105 <220> FEATURE:
106 <221> NAME/KEY: MISC_FEATURE
107 <222> LOCATION: (16)..(17)
108 <223> OTHER INFORMATION: Where the amino acids located at 16 and 17 are hydrophobic
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109      acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala
112 <220> FEATURE:
113 <221> NAME/KEY: MISC_FEATURE
114 <222> LOCATION: (22)..(22)
115 <223> OTHER INFORMATION: Where the amino acid located at 22 is an uncharged polar
amino ac
116      id, including Thr, Ser, Tyr, Gly, Gln, and Asn
119 <220> FEATURE:
120 <221> NAME/KEY: MISC_FEATURE
121 <222> LOCATION: (26)..(27)
122 <223> OTHER INFORMATION: Where the amino acids located at 26 and 27 are hydrophobic
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123      acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala
126 <220> FEATURE:
127 <221> NAME/KEY: MISC_FEATURE
128 <222> LOCATION: (29)..(30)
129 <223> OTHER INFORMATION: Where the amino acids located at 29 and 30 are hydrophobic
amino
130      acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala
133 <220> FEATURE:

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134 <221> NAME/KEY: MISC\_FEATURE .  
135 <222> LOCATION: (33)..(33)  
136 <223> OTHER INFORMATION: Where the amino acid located at 33 is a hydrophobic amino  
acid, i  
137           ncluding Val, Leu, Ile, Met, Pro, Phe, and Ala  
140 <220> FEATURE:

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 142 <222> LOCATION: (35)..(35)  
 143 <223> OTHER INFORMATION: Where the amino acid located at 35 is an uncharged polar  
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 144 id, including Thr, Ser, Tyr, Gly, Gln, and Asn  
 147 <220> FEATURE:  
 148 <221> NAME/KEY: MISC\_FEATURE  
 149 <222> LOCATION: (37)..(38)  
 150 <223> OTHER INFORMATION: Where the amino acids located at 37 and 38 are hydrophobic  
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 151 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 154 <400> SEQUENCE: 3  
 W--> 156 Xaa Xaa Cys Glu Xaa Cys Glu Xaa Xaa Xaa Lys Glu Xaa Xaa Lys Xaa  
 157 1 5 10 15  
 160 Xaa Asp Asn Asn Lys Xaa Glu Lys Glu Xaa Xaa Asp Xaa Xaa Asp Lys  
 161 20 25 30  
 164 Xaa Cys Xaa Lys Xaa Xaa  
 165 35  
 168 <210> SEQ ID NO: 4  
 169 <211> LENGTH: 39  
 170 <212> TYPE: PRT  
 171 <213> ORGANISM: Homo sapiens  
 173 <220> FEATURE:  
 174 <221> NAME/KEY: MISC\_FEATURE  
 175 <222> LOCATION: (1)..(2)  
 176 <223> OTHER INFORMATION: Where the amino acids located at 1 and 2 are hydrophobic  
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 177 ids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
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 181 <221> NAME/KEY: MISC\_FEATURE  
 182 <222> LOCATION: (3)..(3)  
 183 <223> OTHER INFORMATION: Where the amino acid located at 3 is an uncharged polar  
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 184 d, including Thr, Ser, Tyr, Gly, Gln, and Asn  
 187 <220> FEATURE:  
 188 <221> NAME/KEY: MISC\_FEATURE  
 189 <222> LOCATION: (6)..(6)  
 190 <223> OTHER INFORMATION: Where the amino acid located at 6 is a hydrophobic amino  
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 191 cluding Val, Leu, Ile, Met, Pro, Phe, and Ala  
 194 <220> FEATURE:  
 195 <221> NAME/KEY: MISC\_FEATURE  
 196 <222> LOCATION: (9)..(11)  
 197 <223> OTHER INFORMATION: Where the amino acids located at 9-11 are hydrophobic amino  
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 198 , including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 201 <220> FEATURE:  
 202 <221> NAME/KEY: MISC\_FEATURE  
 203 <222> LOCATION: (14)..(14)  
 204 <223> OTHER INFORMATION: Where the amino acid located at 14 is a hydrophobic amino  
 acid, i  
 205 ncluding Val, Leu, Ile, Met, Pro, Phe, and Ala  
 208 <220> FEATURE:

209 <221> NAME/KEY: MISC\_FEATURE .

210 <222> LOCATION: (15)..(15)

211 <223> OTHER INFORMATION: Where the amino acid located at 15 is an uncharged polar amino ac

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212      id, including Thr, Ser, Tyr, Gly, Gln, and Asn
215 <220> FEATURE:
216 <221> NAME/KEY: MISC_FEATURE
217 <222> LOCATION: (17)..(18)
218 <223> OTHER INFORMATION: Where the amino acids located at 17 and 18 are hydrophobic
amino
219      acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala
222 <220> FEATURE:
223 <221> NAME/KEY: MISC_FEATURE
224 <222> LOCATION: (23)..(23)
225 <223> OTHER INFORMATION: Where the amino acid located 23 is an uncharged polar amino
acid,
226      including Thr, Ser, Tyr, Gly, Gln, and Asn
229 <220> FEATURE:
230 <221> NAME/KEY: MISC_FEATURE
231 <222> LOCATION: (27)..(28)
232 <223> OTHER INFORMATION: Where the amino acids located at 27 and 28 are hydrophobic
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233      acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala
236 <220> FEATURE:
237 <221> NAME/KEY: MISC_FEATURE
238 <222> LOCATION: (30)..(31)
239 <223> OTHER INFORMATION: Where the amino acids located at 30 and 31 are hydrophobic
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240      acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala
243 <220> FEATURE:
244 <221> NAME/KEY: MISC_FEATURE
245 <222> LOCATION: (34)..(34)
246 <223> OTHER INFORMATION: Where the amino acid located at 34 is a hydrophobic amino
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247      ncluding Val, Leu, Ile, Met, Pro, Phe, and Ala
250 <220> FEATURE:
251 <221> NAME/KEY: MISC_FEATURE
252 <222> LOCATION: (36)..(36)
253 <223> OTHER INFORMATION: Where the amino acid located at 36 is an uncharged polar
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254      id, including Thr, Ser, Tyr, Gly, Gln, and Asn
257 <220> FEATURE:
258 <221> NAME/KEY: MISC_FEATURE
259 <222> LOCATION: (38)..(39)
260 <223> OTHER INFORMATION: Where the amino acids located at 38 and 39 are hydrophobic
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261      acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala
264 <400> SEQUENCE: 4
W--> 266 Xaa Xaa Xaa Cys Glu Xaa Cys Glu Xaa Xaa Xaa Lys Glu Xaa Xaa Lys
267 1          5          10          15
270 Xaa Xaa Asp Asn Asn Lys Xaa Glu Lys Glu Xaa Xaa Asp Xaa Xaa Asp
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275          35
278 <210> SEQ ID NO: 5
279 <211> LENGTH: 38
280 <212> TYPE: PRT

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281 <213> ORGANISM: Homo sapiens .  
283 <220> FEATURE:  
284 <221> NAME/KEY: MISC\_FEATURE



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Input Set : A:\EP.txt

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285 <222> LOCATION: (1)..(1)  
 286 <223> OTHER INFORMATION: Where the amino acid located at 1 is a hydrophobic amino  
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 287 cluding Val, Leu, Ile, Met, Pro, Phe, and Ala  
 290 <220> FEATURE:  
 291 <221> NAME/KEY: MISC\_FEATURE  
 292 <222> LOCATION: (2)..(2)  
 293 <223> OTHER INFORMATION: Where the amino acid located at 2 is an uncharged polar  
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 294 d, including Thr, Ser, Tyr, Gly, Gln, and Asn  
 297 <220> FEATURE:  
 298 <221> NAME/KEY: MISC\_FEATURE  
 299 <222> LOCATION: (5)..(5)  
 300 <223> OTHER INFORMATION: Where the amino acid located at 5 is a hydrophobic amino  
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 301 cluding Val, Leu, Ile, Met, Pro, Phe, and Ala  
 304 <220> FEATURE:  
 305 <221> NAME/KEY: MISC\_FEATURE  
 306 <222> LOCATION: (8)..(10)  
 307 <223> OTHER INFORMATION: Where the amino acids located at 8-10 are hydrophobic amino  
 acids  
 308 , including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 311 <220> FEATURE:  
 312 <221> NAME/KEY: MISC\_FEATURE  
 313 <222> LOCATION: (13)..(13)  
 314 <223> OTHER INFORMATION: Where the amino acid located at 13 is a hydrophobic amino  
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 315 ncluding Val, Leu, Ile, Met, Pro, Phe, and Ala  
 318 <220> FEATURE:  
 319 <221> NAME/KEY: MISC\_FEATURE  
 320 <222> LOCATION: (14)..(14)  
 321 <223> OTHER INFORMATION: Where the amino acid located at 14 is an uncharged polar  
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 322 id, including Thr, Ser, Tyr, Gly, Gln, and Asn  
 325 <220> FEATURE:  
 326 <221> NAME/KEY: MISC\_FEATURE  
 327 <222> LOCATION: (16)..(17)  
 328 <223> OTHER INFORMATION: Where the amino acids located at 16 and 17 are hydrophobic  
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 329 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 332 <220> FEATURE:  
 333 <221> NAME/KEY: MISC\_FEATURE  
 334 <222> LOCATION: (22)..(22)  
 335 <223> OTHER INFORMATION: Where the amino acid located at 22 is an uncharged polar  
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 336 id, including Thr, Ser, Tyr, Gly, Gln, and Asn  
 339 <220> FEATURE:  
 340 <221> NAME/KEY: MISC\_FEATURE  
 341 <222> LOCATION: (26)..(27)  
 342 <223> OTHER INFORMATION: Where the amino acids located at 26 and 27 are hydrophobic  
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 343 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 346 <220> FEATURE:

347 <221> NAME/KEY: MISC\_FEATURE  
348 <222> LOCATION: (29)..(30)  
349 <223> OTHER INFORMATION: Where the amino acids located at 29 and 30 are hydrophobic  
amino  
350 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
353 <220> FEATURE:

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/780,438C

DATE: 05/08/2003  
TIME: 14:42:58

Input Set : A:\EP.txt  
Output Set: N:\CRF4\05082003\I780438C.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 1,2,5,8,9,10,13,14,16,17,22,26,27,29,30,33,35,37,38  
Seq#:4; Xaa Pos. 1,2,3,6,9,10,11,14,15,17,18,23,27,28,30,31,34,36,38,39  
Seq#:5; Xaa Pos. 1,2,5,8,9,10,13,14,16,17,22,26,27,29,30,33,35,37,38  
Seq#:6; Xaa Pos. 1,2,5,8,9,10,13,14,16,17,22,26,27,29,30,33,35,37,38

## VERIFICATION SUMMARY

DATE: 05/08/2003

PATENT APPLICATION: US/09/780,438C

TIME: 14:42:58

Input Set : A:\EP.txt

Output Set: N:\CRF4\05082003\I780438C.raw

L:156 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
M:341 Repeated in SeqNo=3  
L:266 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0  
M:341 Repeated in SeqNo=4  
L:376 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0  
M:341 Repeated in SeqNo=5  
L:486 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0  
M:341 Repeated in SeqNo=6